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FIG. 1

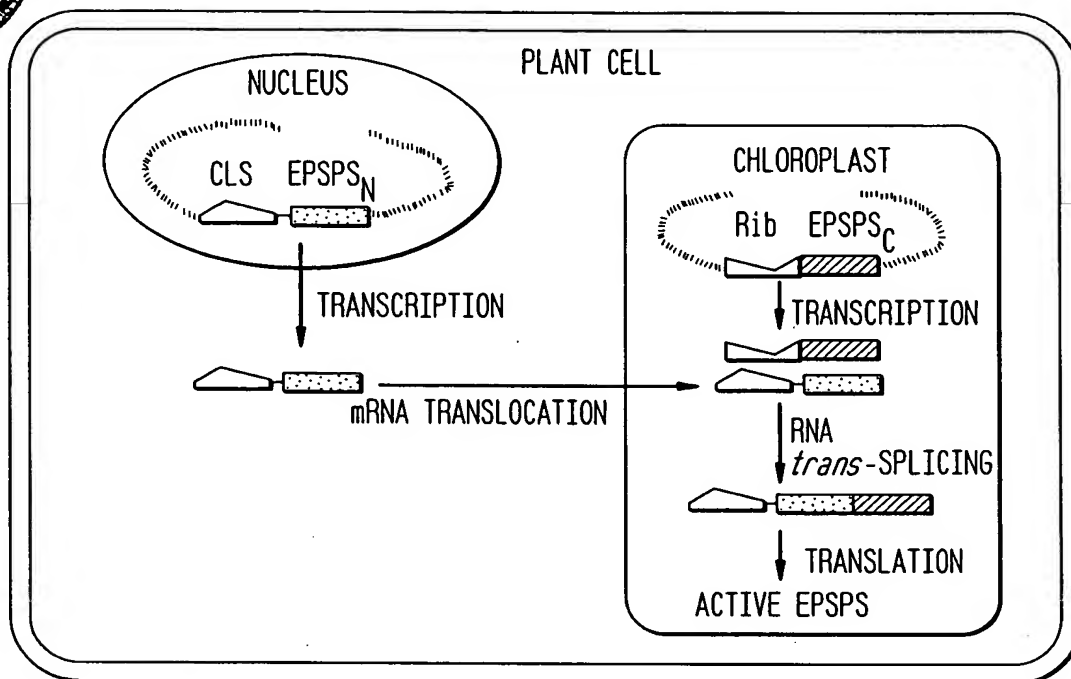
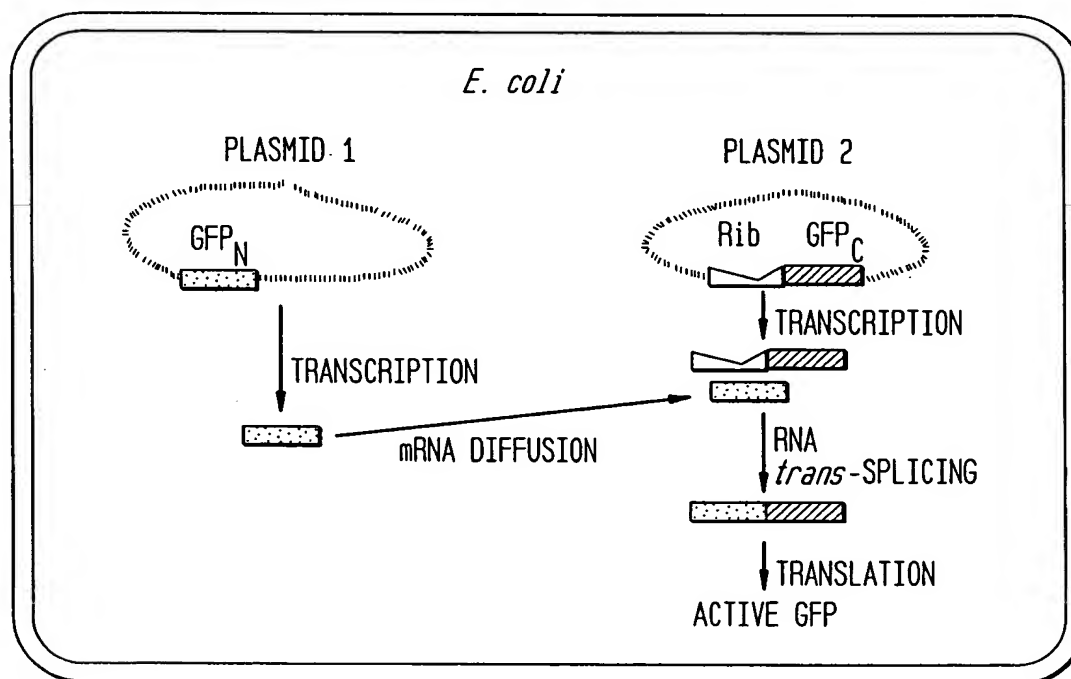


FIG. 2



[illegible]

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FIG. 4

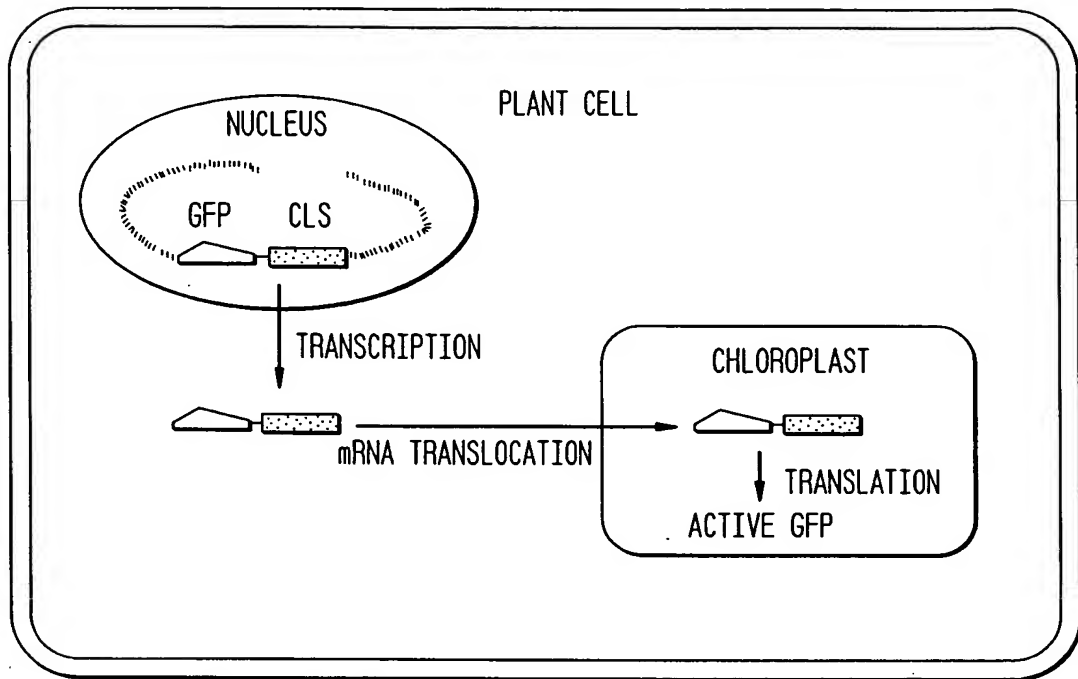
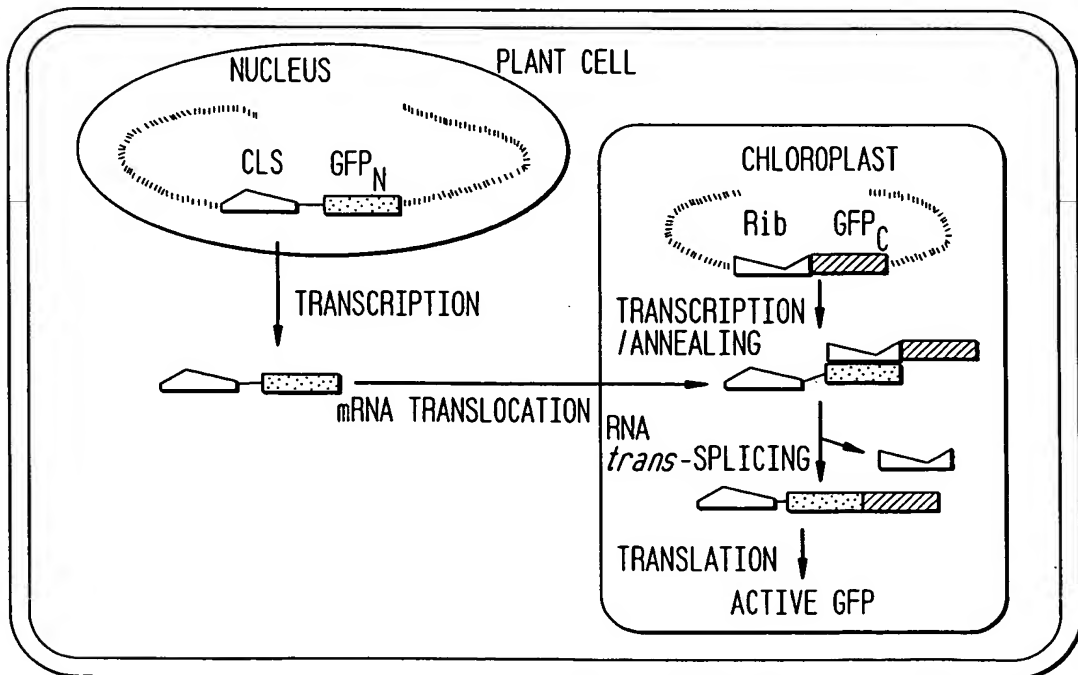


FIG. 5



**FIG. 6**

SEQ ID NO:3

1 tttattagaa caagaagtga ggatatgatt aaactttgtt tgacgaaacc  
51 aggtctgttc cgactttccg actctgagtt tcgacttggt agagaaggag  
101 gagtcgtggt gaacttttat taaaaaaatt agttcactcg tcttcaatct  
151 cttgatcact tcgtctcttc agggaaagat gggaagaaca ctgatgagtc  
201 tcgcaagggt tactcctcta tcttcattgt ttttttacia aatcttg

**FIG. 7**

**FIG. 8A**

pVUG1 segment 5689-6757  
(SEQ ID NO:4)

5689 tttattagaa caagaagtga ggatatgatt aaactttggt tgacgaaacc  
5739 aggtctgttc cgactttccg actctgagtt tcgacttggt agagaaggag  
5789 gagtcgtggt gaacttttat taaaaaaatt agttcactcg tcttcaatct  
5839 cttgatcact tcgtgctagc atgtatttgg caaatcaaat accatggtct  
5889 aataatcaaa cattctgatt agttgataat attagtatta gttggaaatt  
5939 ttgtgaaaga ttcttatgaa aagtttcatt aacacggaat tcgtgtcgag  
5989 tagaccttgt tgttgtgaga attcttaatt catgagttgt agggagggat  
6039 ttATGAGTAA AGGAGAAGAA CTTTTCCTG GAGTTGTCCC AATCTTGT  
6089 GAATTAGATG GTGATGTTAA TGGGCACAAA TTTCTGTCA GTGGAGAGGG  
6139 TGAAGGTGAT GCAACATACG GAAACTTAC CCTTAAATTT ATTTGCACTA  
6189 CTGGAAACT ACCTGTTCCA TGGCCAACAC TTGTCCTAC TTTCTCTAT  
6239 GGTGTTCAAT GCTTTTCAAG ATACCCAGAT CATATGAAGC GGCACGACTT  
6289 CTTCAAGAGC GCCATGCCTG AGGATACGT GCAGGAGAGG ACCATCTCTT  
6339 TCAAGGACGA CGGGAACCTAC AAGACACGTG CTGAAGTCAA GTTTGAGGGA  
6389 GACACCCTCG TCAACAGGAT CGAGCTTAAG GGAATCGATT TCAAGGAGGA  
6439 CGGAAACATC CTCGGCCACA AGTTGGAATA CAACTACAAC TCCCACAACG  
6489 TATACATCAC GGCAGACAAA CAAAAGAATG GAATCAAAGC TAACTTCAAA  
6539 ATTAGACACA ACATTGAAGA TGAAGCGTT CAACTAGCAG ACCATTATCA  
6589 ACAAATACT CCAATTGGCG ATGGCCCTGT CCTTTTACCA GACAACCATT  
6639 ACCTGTCCAC ACAATCTGCC CTTTCGAAAG ATCCCAACGA AAAGAGAGAC  
6689 CACATGGTCC TTCTTGAGTT TGTAACAGCT GCTGGGATTA CACATGGCAT  
6739 GGATGAACTA TACAAATAA

**FIG. 8B**

pVUG2 segment 5689-6757  
(SEQ ID NO:5)

5689 acttgtgaga gaaggaggag tcgtggtgaa cttttattaa aaaaattagt  
5739 tcactcgtct tcaatctctt gatcacttcg tctcttcagg gaaagatggg  
5789 aagaacactg atgagtctcg caaggtttac tcctctatct tcattgtttt  
5839 ttacaaaaat cttggctagc atgtatttgg caaatcaaat accatggctt  
5889 aataatcaaa cattctgatt agttgataat attagtatta gttggaaatt  
5939 ttgtgaaaga ttcctatgaa aagtttcatt aacacggaat tcgtgtcgag  
5989 tagaccttgt tgttgtgaga attcttaatt catgagttgt agggagggat  
6039 ttATGAGTAA AGGAGAAGAA CTTTCACTG GAGTTGTCCC AATTCTTGT  
6089 GAATTAGATG GTGATGTTAA TGGGCACAAA TTTTCTGTCA GTGGAGAGGG  
6139 TGAAGGTGAT GCAACATACG GAAACTTAC CCTTAAATTT ATTTGCACTA  
6189 CTGGAAACT ACCTGTTCCA TGGCCAACAC TTGTCCTAC TTTCTCTTAT  
6239 GGTGTTCAAT GCTTTTCAAG ATACCCAGAT CATATGAAGC GGCACGACTT  
6289 CTTCAAGAGC GCCATGCCTG AGGGATACGT GCAGGAGAGG ACCATCTCTT  
6339 TCAAGGACGA CGGGAACACT AAGACACGTG CTGAAGTCAA GTTTGAGGGA  
6389 GACACCCTCG TCAACAGGAT CGAGCTTAAG GGAATCGATT TCAAGGAGGA  
6439 CGGAAACATC CTCGGCCACA AGTTGGAATA CAACTACAAC TCCCACAACG  
6489 TATACATCAC GGCAGACAAA CAAAAGAATG GAATCAAAGC TAACTTCAAA  
6539 ATTAGACACA ACATTGAAGA TGGAAGCGTT CAACTAGCAG ACCATTATCA  
6589 ACAAATACT CCAATTGGCG ATGGCCCTGT CCTTTTACCA GACAACCATT  
6639 ACCTGTCCAC ACAATCTGCC CTTTCGAAAG ATCCCAACGA AAAGAGAGAC  
6689 CACATGGTCC TTCTTGAGTT TGTAACAGCT GCTGGGATTA CACATGGCAT  
6739 GGATGAACTA TACAAATAA

**FIG. 8C**

pVUG3 segment 6833-7901  
(SEQ ID NO:6)

6833 tcttcagga aagatggga gaacactgat gagtctcgca aggtttactc  
6883 ctctatcttc attgtttttt tacaaaaatct tgtttattag aacaagaagt  
6933 gaggatatga ttaaactttg ttgacgaaa ccaggctgt tccgactttc  
6983 cgactctgag tttcgctagc atgtatttgg caaatcaaat accatggtct  
7033 aataatcaaa cattctgatt agttgataat attagtatta gttggaaatt  
7083 ttgtgaaaga ttcctatgaa aagtttcatt aacacggaat tcgtgtcgag  
7133 tagaccttgt tgttgtgaga attcttaatt catgagttgt agggagggat  
7183 ttATGAGTAA AGGAGAAGAA CTTTCTACTG GAGTTGTCCC AATTCTTGTT  
7233 GAATTAGATG GTGATGTTAA TGGGCACAAA TTTCTGTCA GTGGAGAGGG  
7283 TGAAGGTGAT GCAACATACG GAAACTTAC CCTTAAATTT ATTTGCACTA  
7333 CTGGAAACT ACCTGTTCCA TGGCCAACAC TTGTCACTAC TTTCTCTTAT  
7383 GGTGTTCAAT GCTTTTCAAG ATACCCAGAT CATATGAAGC GGCACGACTT  
7433 CTTCAAGAGC GCCATGCCTG AGGGATACGT GCAGGAGAGG ACCATCTCTT  
7483 TCAAGGACGA CGGGAACAC AAGACACGTG CTGAAGTCAA GTTTGAGGGA  
7533 GACACCCTCG TCAACAGGAT CGAGCTTAAG GGAATCGATT TCAAGGAGGA  
7583 CGGAAACATC CTCGGCCACA AGTTGGAATA CAACTACAAC TCCCACAACG  
7633 TATACATCAC GGCAGACAAA CAAAAGAATG GAATCAAAGC TAACTTCAAA  
7683 ATTAGACACA ACATTGAAGA TGGAAGCGTT CAACTAGCAG ACCATTATCA  
7733 ACAAATACT CCAATTGGCG ATGGCCCTGT CCTTTTACCA GACAACCATT  
7783 ACCTGTCCAC ACAATCTGCC CTTTCGAAAG ATCCCAACGA AAAGAGAGAC  
7833 CACATGGTCC TTCTTGAGTT TGTAACAGCT GCTGGGATTA CACATGGCAT  
7883 GGATGAACTA TACAAATAA

**FIG. 9**

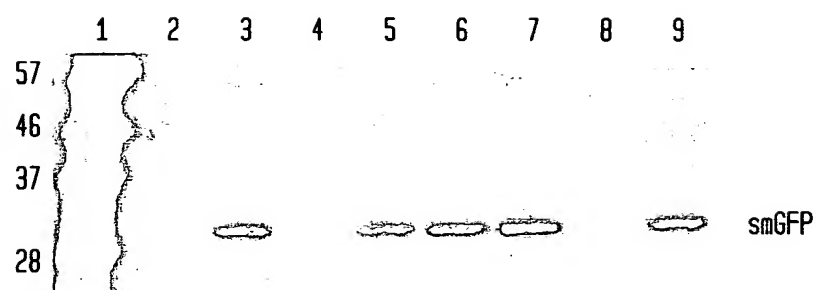




FIG. 10

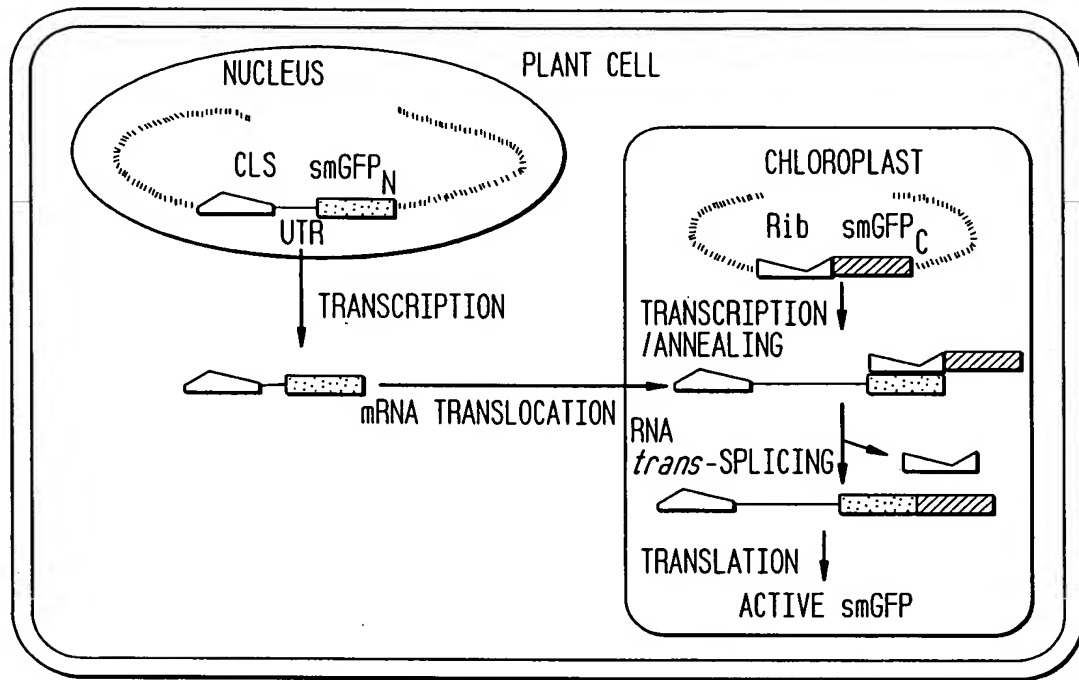


FIG. 11A

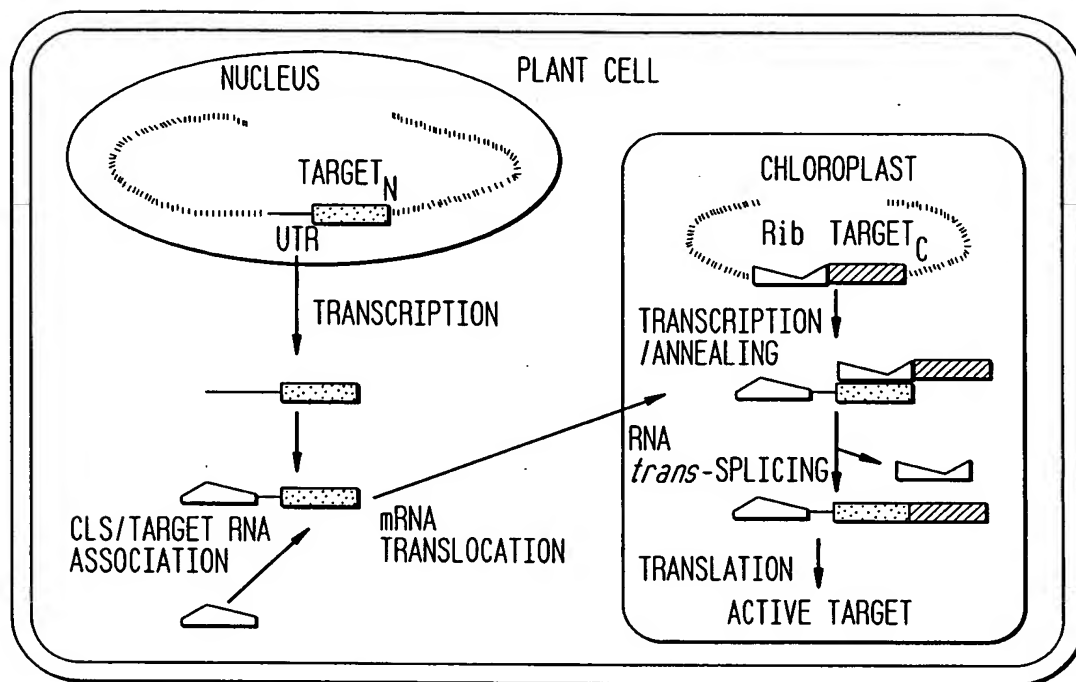


FIG. 11B

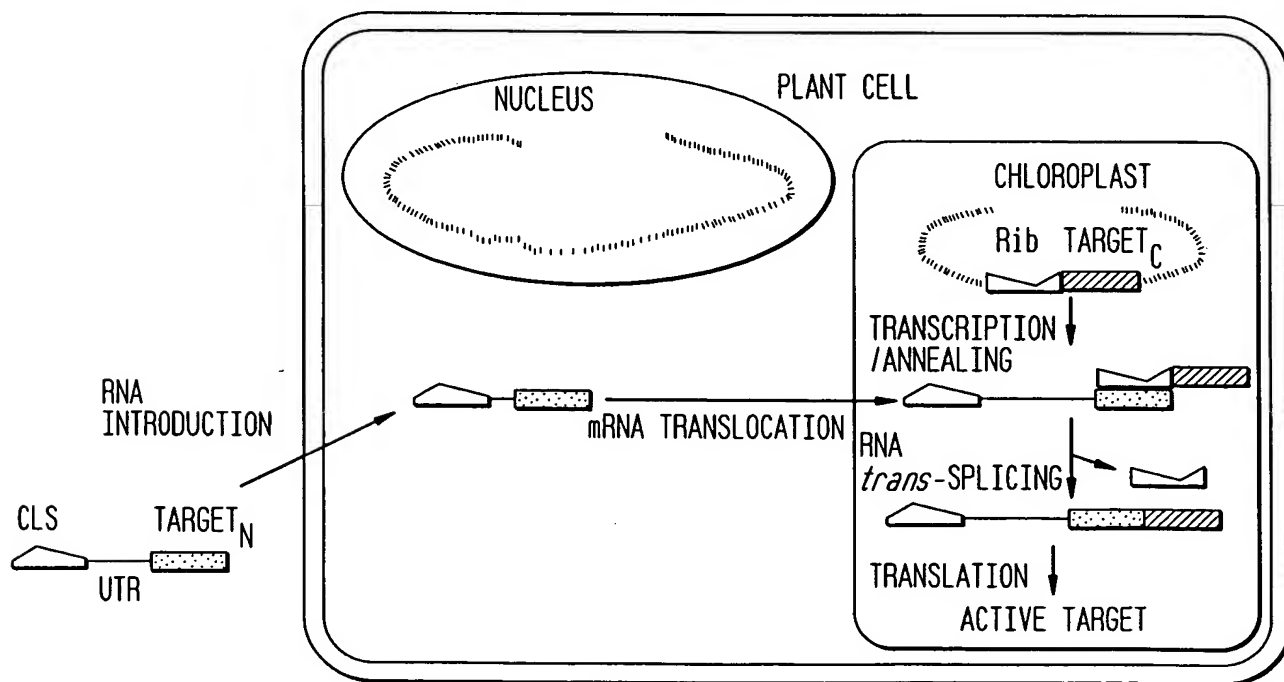


FIG. 11C

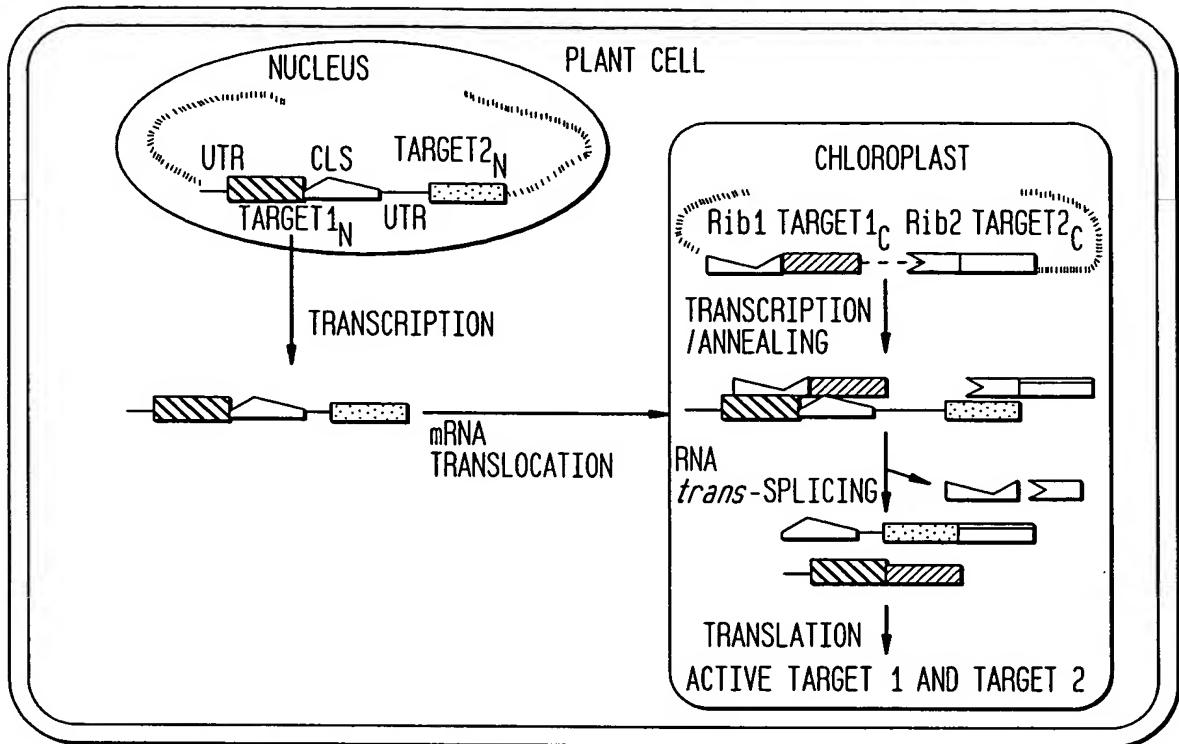


FIG. 11D

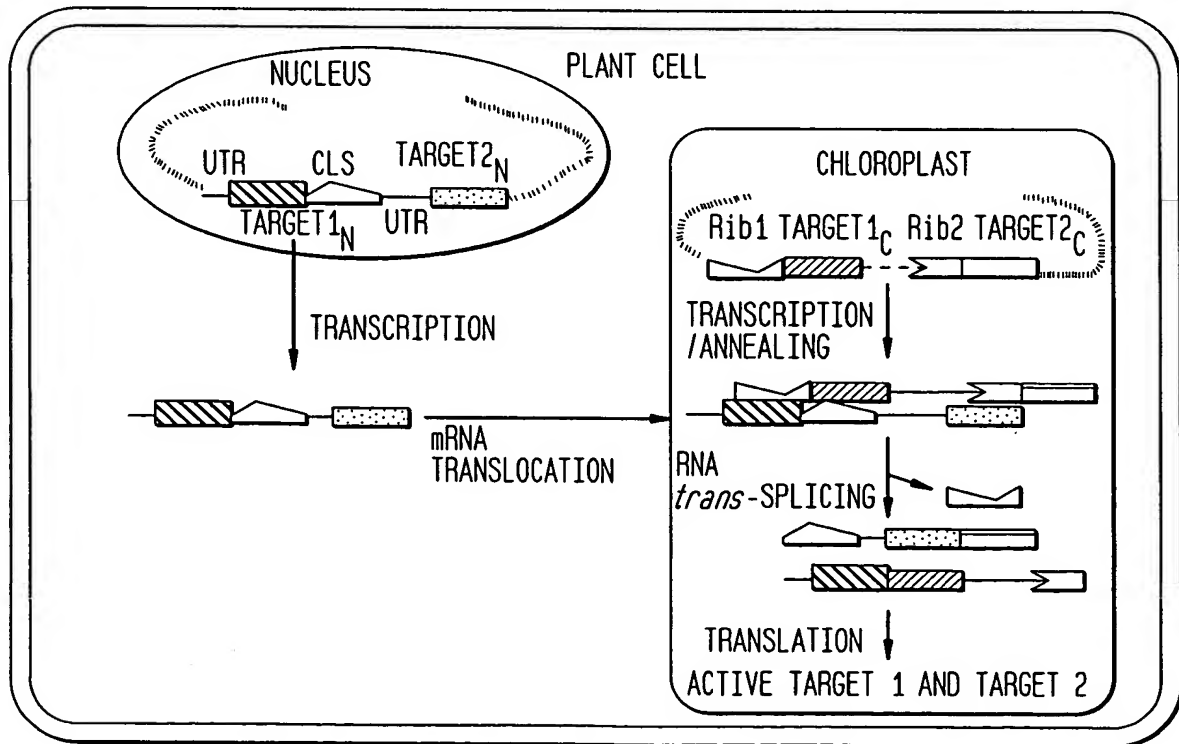


FIG. 11E

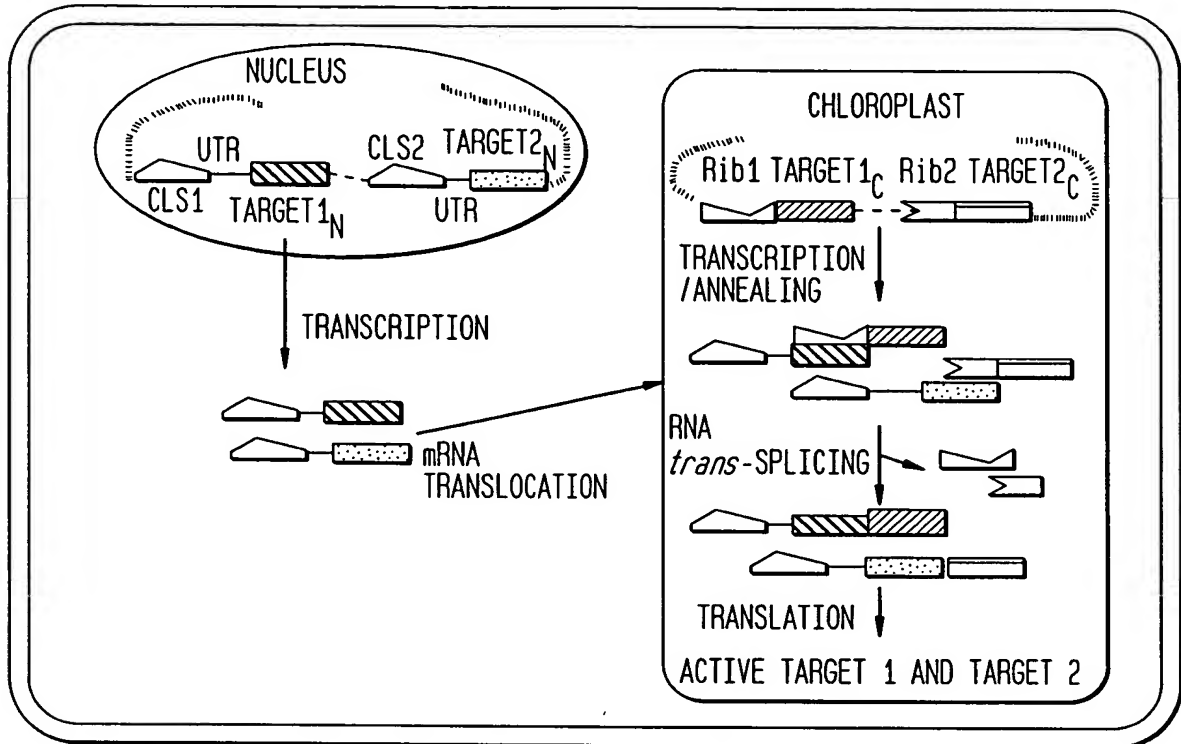


FIG. 11F

